

# UC Davis

## UC Davis Previously Published Works

### Title

Corrigendum: Metagenomic and whole-genome analysis reveals new lineages of gokushoviruses and biogeographic separation in the sea.

### Permalink

<https://escholarship.org/uc/item/2415v3nv>

### Authors

Labonté, Jessica M  
Suttle, Curtis A

### Publication Date

2015

### DOI

10.3389/fmicb.2015.00114

Peer reviewed



# Corrigendum: Metagenomic and whole-genome analysis reveals new lineages of gokushoviruses and biogeographic separation in the sea

Jessica M. Labonté<sup>1†</sup> and Curtis A. Suttle<sup>1,2,3,4\*</sup>

<sup>1</sup> Department of Microbiology and Immunology, University of British Columbia, Vancouver, BC, Canada

<sup>2</sup> Department of Earth, Ocean and Atmospheric Sciences, University of British Columbia, Vancouver, BC, Canada

<sup>3</sup> Department of Botany, University of British Columbia, Vancouver, BC, Canada

<sup>4</sup> Integrated Microbial Diversity, Canadian Institute for Advanced Research, University of British Columbia, Vancouver, BC, Canada

\*Correspondence: [suttle@science.ubc.ca](mailto:suttle@science.ubc.ca)

<sup>†</sup> Present address:

Jessica M. Labonté, Bigelow Laboratory for Ocean Sciences, East Boothbay, USA

Edited and reviewed by:

Brian Palenik, Scripps Institution of Oceanography, USA

**Keywords:** biogeography, ssDNA viruses, *Microviridae*, *Gokushovirinae*, virus diversity, ocean viruses

## A corrigendum on

Metagenomic and whole-genome analysis reveals new lineages of gokushoviruses and biogeographic separation in the sea by Labonté, J. M., and Suttle, C. A. (2013). *Front. Microbiol.* 4:404. doi: 10.3389/fmicb.2013.00404

The primer MicroVP1-R1 should read as “5′-NCG YTC YTG RTA NCC RAA-3′.”

**Conflict of Interest Statement:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Received: 26 January 2015; accepted: 29 January 2015; published online: 11 February 2015.

Citation: Labonté JM and Suttle CA (2015) Corrigendum: Metagenomic and whole-genome analysis reveals new lineages of gokushoviruses and biogeographic separation in the sea. *Front. Microbiol.* 6:114. doi: 10.3389/fmicb.2015.00114

This article was submitted to *Evolutionary and Genomic Microbiology*, a section of the journal *Frontiers in Microbiology*.

Copyright © 2015 Labonté and Suttle. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) or licensor are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.